

Functional Diversity of Histamine and Histamine Receptors

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In order to analyze the mechanisms by which a single biogenic amine like histamine is capable of inducing a wide variety of both physiologic and pathologic functions in various tissues/cells, histamine responses were dissected in detail from a biochemical and pharmacologic point of view. Histamine is synthesized by multiple isozymes of histidine decarboxylase, and catabolized by either diamine oxidase or histamine-N-methyltransferase. Synthesized intracellular histamine may play a role in cell proliferation, whereas released histamine binds to at least three different histamine-

specific receptors, then activates various intracellular components, such as Ca^{++} , cAMP, protein kinase, and ion channels. These second messenger pathways interact differentially with each other in various tissues/cells. Moreover, histamine not only activates its own receptors, but also activates other related receptors such as the serotonin 1c receptor. Therefore, to understand the complex actions of histamine, new approaches should be established, in which multiple phenomena can be monitored simultaneously. *J Invest Dermatol* 98:8S-11S, 1992

Histamine is known to exhibit a wide variety of both physiologic and pathologic functions in different tissues and cells, and also appears to serve as a neurotransmitter/neuromodulator in the brain [1]. Histamine action is believed to be mediated by binding to its specific cell-surface receptors. Extensive pharmacologic analysis identifies at least three different subtypes of histamine receptors using highly specific competitive agonists and antagonists. The H_1 histamine receptor is mainly involved in smooth muscle contraction and vascular permeability [2]. The H_2 receptor mediates gastric acid secretion [3], whereas the H_3 receptor is present at pre-synaptic locations in the central nervous system and inhibits the release of histamine from depolarized slices of rat brain [4].

Although three histamine receptors have been studied exten-

sively from the pharmacologic point of view [5,6], the biochemical nature of histamine receptors are incompletely characterized. Furthermore, growing evidence suggests that complex and heterogeneous activities of histamine in vitro and in vivo cannot be explained by three subtypes of histamine receptors alone. Therefore the purpose of this article is to delineate the heterogeneous actions of histamine biochemically, and to understand how a single biogenic amine is capable of inducing a variety of different cellular functions under different circumstances.

Heterogeneity of Histamine Metabolism Histamine in tissues is derived from the decarboxylation of histidine in a single step by histidine decarboxylase (HDC) (Fig 1). A number of studies suggest that HDC is not a homogenous enzyme, in which multiple isoforms of HDC exhibitions with different isoelectric points have been reported [7,8]. This heterogeneity of HDC is further supported by the evidence that at least two different HDC cDNA have been cloned that show 86.1% homology in their amino acid sequences [9,10]. Furthermore, histamine is also derived from histidine by aromatic-L-amino acid decarboxylase (DOPA decarboxylase, DDC) [11]. Amino acid sequence analysis suggests that the NH_2 -terminal 483 amino acid residues of the cloned rat HDC [9] shared 52% and 53% identities with those of rat DDC [12] and guinea-pig DDC [13], respectively.

Histamine is released in conjunction with other potent chemical mediators in response to specific IgE-mediated [14] or non-specific mechanisms [15] from the tissue mast cells [16] or blood basophilic leukocytes [17]. Released histamine is then catabolized by either diamine oxidase (DAO) or histamine-N-methyltransferase (HMT) [11] (Fig 1). The dominant pathways are different in various species and in different organs/tissues. For example, HMT is the major pathway in mammalian brain, whereas DAO is the main metabolic pathways in the nervous system of some invertebrates [11].

Heterogeneity of H_1 Receptors An early demonstration of the presence of the H_1 receptor on the membrane fractions of guinea pig ileum smooth muscle was carried out by Hill et al [18] using the radiolabeled H_1 -specific antagonist [^3H]-pyrilamine. The dissocia-

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Abbreviations:

AC: adenylyl cyclase
cAMP: cyclic adenosine monophosphate
DAO: diamine oxidase
DDC: DOPA decarboxylase
DPPE: N,N-diethyl-2-[4-(phenylmethyl)phenoxy]-ethanamine-HCl
 EC_{50} : a half maximal concentration
HDC: histidine decarboxylase
HMT: histamine-N-methyltransferase
HPLC: high-performance liquid chromatography
 IP_3 : inositol trisphosphate
kd: dissociation constant
PCR: polymerase chain reaction
 PIP_2 : phosphatidylinositol diphosphate
PKC: protein kinase C
PLC: phospholipase C
SDS-PAGE: sodium dodecyl sulfate polyacrylamide gel electrophoresis
SHT: 5-hydroxytryptamine

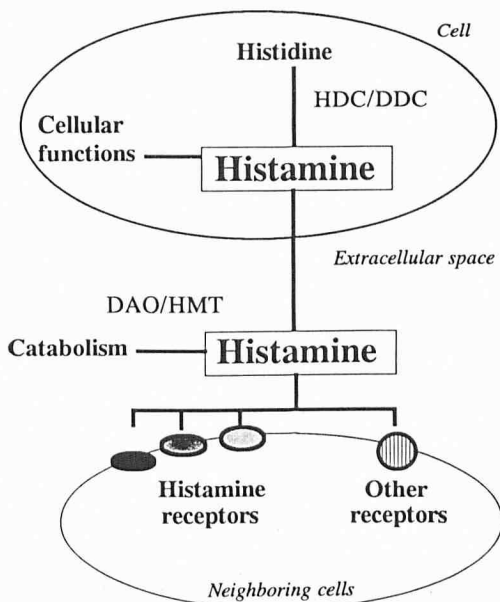


Figure 1. Heterogenous activities of histamine. Histamine is synthesized by histidine decarboxylase (HDC) or DOPA decarboxylase (DDC), and catabolized by either diamine oxidase (DAO) or histamine-N-methyltransferase (HMT). Synthesized intracellular histamine may induce cellular functions such as cell proliferation and platelet aggregation, whereas released histamine binds to at least three different histamine specific receptors (H_1 , H_2 , and H_3 receptors), and is also capable of activating other related receptors like serotonin 1c receptors.

tion constant (K_d) from [3H]-pyrilamine binding studies correlated with the EC_{50} of the contractile response of guinea pig ileum to histamine, suggesting that [3H]-pyrilamine binding sites were the specific H_1 receptor. However, a similar number of [3H]-pyrilamine binding sites were demonstrated on rat, guinea pig, and rabbit ileal membranes, although the contractile effects of histamine varied widely in these species [19]. The K_d of [3H]-pyrilamine also varied widely, from 0.7 nM to 219 nM in various tissues [20]. The difference of activity of H_1 antagonists was also demonstrated in functional assays, such as smooth muscle contraction, Ca^{++} mobilization, glycogen hydrolysis, prostaglandin E synthesis, ion transport, Ca^{++} efflux, and cGMP formation, with an IC_{50} ranging from 30 pM to 10 μ M [20]. Moreover, subclasses of the H_1 receptor were detected in various tissues using different radiolabeled ligands [20].

[3H]-pyrilamine binding studies indicate that extracellular Na^+ , Mg^{++} , Mn^{++} , and guanine nucleotides selectively alter the agonist affinity in guinea pig brain membranes [21], and dithiothreitol also increased agonist affinity in guinea pig ileum [22]. This might suggest that the heterogeneity of the H_1 receptors is due to the microenvironments in which the H_1 receptor is present, or the additional interaction with G-proteins, as demonstrated with other G-protein-coupled membrane receptors [23].

In our previous studies, we have shown that during cellular differentiation of cultured BC3H1 smooth muscle cells, the K_d and the receptor number for [3H]-pyrilamine binding were transiently decreased from 276 nM to 46.5 nM, and from 13.3×10^6 /cell to 2×10^6 /cell, respectively [24]. Data from saturation binding experiments indicate that the high-affinity H_1 receptor on differentiated cells was converted to the low-affinity receptor by the incubation of BC3H1 cells with glycosylation inhibitors, tunicamycin, or swainsonine without a significant change in receptor number [24]. Tunicamycin and swainsonine had no effect on the K_d of the low-affinity H_1 receptor on undifferentiated cells. Concanavalin A, wheat germ agglutinin, lentil lectin, and phytohemagglutinin, which bind to carbohydrate molecules, also inhibited [3H]-pyrilamine binding on differentiated cell membranes, but had no effect on undifferentiated cell membranes [24]. The molecular weight of the H_1 receptor on

differentiated cells was approximately 68,000 as analyzed by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), which after treatment with N-glycanase was shifted to 40,000, a molecular weight similar to that of the H_1 receptor on undifferentiated cells [24]. These results suggest that one element contributing to the H_1 -receptor heterogeneity is receptor N-glycosylation.

Ruat et al [25] have used irreversible photoaffinity labeling to specifically label H_1 -binding proteins in cell membranes. They synthesized [^{125}I]-iodoazidophenpyramine, a highly potent H_1 receptor antagonist derived from pyrilamine. Upon irradiation, 5% of the bound radioactivity was covalently incorporated into the H_1 receptor of guinea pig brain membranes. SDS-PAGE analysis indicated that the molecular weight of the H_1 receptor was 350,000–400,000 dalton in the absence of 2-mercaptoethanol, and the molecular weight decreased to 56,000 and 47,000 in the presence of 2-mercaptoethanol. This suggests that the H_1 receptor may have one or more disulfide bridges, resulting in a higher molecular weight complex. Furthermore, in the presence of protease inhibitors, labeling of the 56,000-dalton peptide increased at the expense of the 47,000-dalton peptide. This also suggests that the 47,000-dalton peptide is a proteolysis product of the 56,000-dalton peptide. In our previous studies, the protein with [3H]-pyrilamine binding activity was purified with gel filtration, chromatofocusing followed by reverse-phase high-performance liquid chromatography (HPLC) from hamster vas deferens DDT1MF-2 cells [26]. The final material was a single peak on reverse-phase HPLC and a single band of apparent molecular weight 39,000 daltons on SDS-PAGE, which exhibited specific [3H]-pyrilamine binding activity. This molecular weight of the H_1 receptor on DDT1MF-2 is smaller than that reported by Ruat et al [25], and might also be due to proteolytic degradation. Furthermore, as described above [24], part of the difference of molecular weight might also be due to receptor glycosylation.

The H_1 receptor is known to be coupled to phosphatidylinositol hydrolysis pathways, in which the H_1 receptor-specific stimulation induced hydrolysis of phosphatidylinositol with formation of inositol-1,4,5-trisphosphate (IP_3) and diacylglycerol, leading to intracellular Ca^{++} mobilization and activation of protein kinase C (PKC) [1]. Recent molecular cloning studies have identified multiple subtypes of PKC [27]. Therefore, action of PKC after stimulation with the H_1 receptor may be varied in different cells depending upon the cellular components of PKC subtypes (Fig 2).

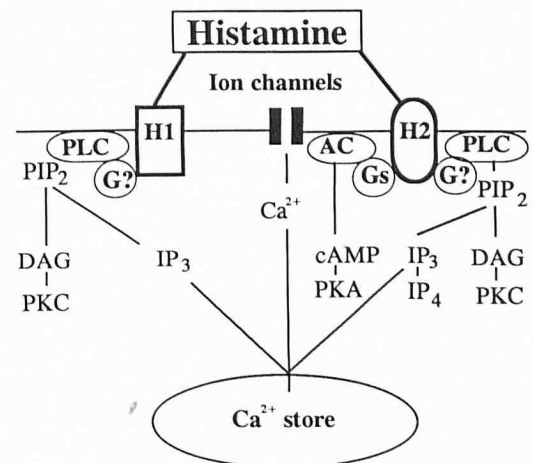


Figure 2. Functional diversity of the H_1 and H_2 receptors. The H_1 receptor is coupled to unidentified G proteins ($G?$), which activate phospholipase C (PLC) followed by hydrolysis of phosphatidylinositol diphosphate (PIP_2) to 1,4,5-inositol trisphosphate (IP_3) and diacylglycerol (DAG). IP_3 induces Ca^{++} mobilization from intracellular stores, and DAG is known to activate protein kinase C (PKC). The H_2 receptor is capable of activating both adenylyl cyclase (AC) and PLC by coupling to two different G proteins (Gs and $G?$).

A number of phosphatidylinositol-coupled Ca^{++} -mobilizing receptors have been already cloned, and sequence homology analysis indicates the striking homology among these receptor cDNA. Therefore, the H_1 receptor cDNA will be cloned by a low-stringency hybridization technique or polymerase chain reaction (PCR) in the near future.

Functional Heterogeneity of H_2 Receptors As described above, H_1 receptors are known to be coupled to phosphatidylinositol hydrolysis pathways, whereas the H_2 receptor is coupled to adenylate cyclase with histamine-stimulating increases in intracellular cyclic adenosine monophosphate (cAMP) [3]. Both the H_1 and H_2 receptors are co-localized in various cells/tissues, and regulate histamine responses via secondary Ca^{++} and cAMP pathways, respectively [28].

In neutrophils, H_2 -specific stimulation consistently enhances the chemokinetic response, whereas the chemotactic response was inhibited in a H_2 -receptor-dependent manner [29,30]. Although the inhibition of chemotaxis is due to the elevation of cAMP, the mechanism of the H_2 -receptor-mediated stimulation of chemokinesis is not explained by the activation of secondary cAMP pathways. This suggests that the H_2 receptor could have multiple independent effects on steps early in neutrophil activation.

Chew reported that the H_2 -receptor stimulation not only activated cAMP pathways, but also increased $[\text{Ca}^{++}]_i$ in parietal cells [31]. In our previous studies, we have also shown that histamine increased $[\text{Ca}^{++}]_i$ in human HL-60 granulocytic cells in a dose-dependent manner with a half-maximal concentration (EC_{50}) of approximately 10^{-6} – 10^{-5} M, which exhibited H_2 -receptor specificity [32]. The cell-permeable form of cAMP and forskolin failed to increase $[\text{Ca}^{++}]_i$ [32]. Because the H_2 agonist, 4-methyl histamine-induced mobilization of $[\text{Ca}^{++}]_i$ was not diminished by reducing the concentration of extracellular Ca^{++} nor by the addition of Ca^{++} channel antagonists, LaCl_3 and nifedipine, we found $[\text{Ca}^{++}]_i$ mobilization was due to the release of Ca^{++} from intracellular stores. Furthermore, histamine increased the levels of IP_3 [32], suggesting that the H_2 receptor is capable of activating phosphatidylinositol hydrolysis pathways like the H_1 receptor. Interestingly, histamine-induced Ca^{++} mobilization was inhibited by the preincubation with cholera toxin, but not by pertussis toxin. Cholera toxin is known to irreversibly activate Gs protein, resulting in increasing intracellular cAMP. Therefore, the H_2 receptor on HL-60 cells may be coupled to two different cholera-toxin-sensitive G-proteins, and activate adenylate cyclase and phospholipase C simultaneously (Fig 2).

The H_2 receptor-mediated intracellular signaling pathways can be altered by modification of the ligand. For example, histamine albumin conjugates, which were used to identify the H_2 receptor-bearing leukocytes for many years [33], have been shown to act as active ligands and increase the level of $[\text{Ca}^{++}]_i$ in HL-60 cells in an H_2 -receptor-specific manner [34]. Both histamine and histamine-albumin conjugates specifically desensitized each other, but the chemotactic peptide fMLP failed to interact with both ligands, although histamine and fMLP increased $[\text{Ca}^{++}]_i$ to a similar degree in HL-60 cells [34]. However, unlike histamine, histamine-albumin-induced $[\text{Ca}^{++}]_i$ mobilization was sustained for more than 10 min as a result of Ca^{++} influx [34]. The H_2 receptor has been shown to interact with Ca^{++} channels as a result of cAMP-dependent phosphorylation of Ca^{++} channels in other systems [35]. However, in HL-60 cells, the H_2 receptor-specific stimulation failed to activate $^{45}\text{Ca}^{++}$ influx, although histamine has been shown to increase cAMP in the H_2 -receptor-specific manner [36]. More interestingly, the level of 1,3,4,5- IP_4 was also high and sustained after stimulation with histamine-albumin conjugates [37]. As IP_4 has been shown to activate Ca^{++} channels [38], histamine-albumin-induced Ca^{++} influx may be due to the altered metabolism of IP_3 , rather than to cAMP-mediated phosphorylation of Ca^{++} channels.

Recently, the cDNA for the H_2 receptor has been cloned using a PCR approach from gastric parietal cells [39]. Its amino acid sequence is shorter than other G-protein-coupled receptor sequences, but exhibits similar structural motifs, in which the peptide spans the membrane seven times with an extracellular NH_2 -termi-

nal and intracellular COOH-terminal. Furthermore, the first extracellular domain contains a possible N-glycosylation site, and cysteine residues are found in both the second and the third extracellular domains [39]. However, Northern blot analysis shows that mRNA of the H_2 receptor is not detectable in heart, whereas the H_2 receptor mediates a positive chronotropic effect in right atria [40] and a positive inotropic effect in ventricle via cAMP [41]. As recent gene cloning techniques have identified more isoforms of receptor genes than would have been expected from the known pharmacology of those receptors, such as muscarinic acetylcholine receptors [42], GABA receptors [43], and glycine receptors [44], more subtypes of the H_2 receptor may be identified in the future.

Cross-talk Between Histamine and Serotonin Receptors

Serotonin (5-hydroxytryptamine, 5HT) and histamine are biogenic amines with similar structure and similar biologic activities. Extensive pharmacologic studies using various synthetic agonists and antagonists have demonstrated that both 5HT and histamine exert physiologic and pathologic activities through binding to their specific cell-surface receptors. However, it is not well characterized whether 5HT and histamine react only with their own specific receptors, or whether under certain circumstances these ligands functionally cross-react with each other. Such analyses are difficult because most cells/tissues express various subtypes of 5HT and histamine receptors on their cell surface.

Although the H_1 receptor cDNA has not been cloned, three serotonin receptor cDNA (5HT1a, 5HT1c, and 5HT2b) have been cloned using a unique *Xenopus* oocyte expression system [45] and low-stringency cross-hybridization techniques [46,47]. Because the 5HT1c and the H_1 receptors are known to share common intracellular phosphatidylinositol hydrolysis pathways, a system that allows the functional expression of 5HT1c cDNA is a good model for the further analysis of cross-reactivity between 5HT and histamine. In our previous study [48], the RNA for the 5HT1c receptor was transcribed in vitro from the cloned 5HT1c receptor cDNA, and was injected into *Xenopus* oocytes in order to study 5HT1c receptor responses to various biogenic amines. As a result, histamine and the H_1 receptor agonists, but not the H_2 and H_3 agonists, significantly induced $^{45}\text{Ca}^{++}$ efflux in 5HT1c-receptor-RNA-injected oocytes, but not in uninjected and water-injected oocytes. However, the H_1 , H_2 , and H_3 antagonists failed to inhibit histamine-induced $^{45}\text{Ca}^{++}$ efflux at 10^{-6} M. This clearly suggests that the 5HT1c receptor can be stimulated by histamine, although its action is different from "classical" histamine pharmacology.

Histamine as an Intracellular Messenger Action of released histamine was believed to be mediated by various subtypes of histamine receptors as described above. However, because HDC activity and intracellular concentrations of histamine have been correlated with cell growth [49], intracellularly synthesized histamine may have a role in cell growth and tissue repair. More recently, Saxena et al [50] reported that inhibitors of HDC suppressed phorbol ester- or collagen-induced platelet aggregation, which was reversed by the novel histamine non- H_1 , non- H_2 , non- H_3 antagonist, N,N-diethyl-2-[4-(phenylmethyl)phenoxy]-ethanamine-HCl (DPPE). Furthermore, in saponin-permeabilized platelets, added histamine reversed the inhibition by DPPE or HDC inhibitors on platelet aggregation. These data suggest a role for histamine as an intracellular messenger, although it is still unclear whether histamine reacts with intracellular receptors.

CONCLUSION

Histamine is capable of inducing a wide variety of cellular functions through its heterogenous receptors, or by direct action in the cytosol by interacting with other multiple factors. In order to understand the complex actions of histamine in biology and medicine, more studies are required, and perhaps new approaches should be established, in which multiple phenomena can be monitored simultaneously.

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